RAW SEQUENCE LISTING
ERROR REPORT

BIC   :HNOLOGY
SYSTEMS
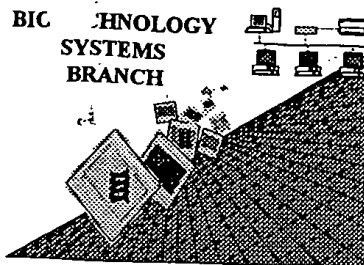BRANCH

A.5

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: **09/868,885**

Source: **Pu/09**

Date Processed by STIC: **7/9/2001**

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: **patin21help@uspto.gov** or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: **patin3help@uspto.gov** or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 3.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:
http://www.uspto.gov/web/offices/pac/checker

Raw Sequence Listing Error Summary

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: 09/868,885

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1 ____ Wrapped Nucleics
Wrapped Aminos · The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 ____ Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 ____ Misaligned Amino Numbering The numbering under each 5ᵗʰ amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4 ____ Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5 ____ Variable Length Sequence(s) ____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 ____ PatentIn 2.0 "bug" · A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7 ____ Skipped Sequences (OLD RULES) Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8 ____ Skipped Sequences (NEW RULES) Sequence(s) ____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 ____ Use of n's or Xaa's (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 ____ Invalid <213> Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 ____ Use of <220> Sequence(s) 44-45 missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12 ____ PatentIn 2.0 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

AMC – Biotechnology Systems Branch – 06/04/2001

PCT

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/868,885

DATE: 07/09/2001
TIME: 10:11:18

Input Set : A:\2584WOOP.txt.txt
Output Set: N:\CRF3\07092001\I868885.raw

Does Not Comply
Corrected Diskette Needed

P-6

```
           3 <110> APPLICANT: Takeda Chemical Industries, Ltd.
W-->       4 <120> TITLE OF INVENTION: Use of Peptide
W-->       5 <130> FILE REFERENCE: 2584WOOP
C-->       6 <140> CURRENT APPLICATION NUMBER: US/09/868,885
C-->       6 <141> CURRENT FILING DATE: 2001-06-22
           6 <150> PRIOR APPLICATION NUMBER: JP 10-369585
           7 <151> PRIOR FILING DATE: 1998-12-25
W-->       8 <160> NUMBER OF SEQ ID: 45
          11 <210> SEQ ID NO: 1
          12 <211> LENGTH: 98
          13 <212> TYPE: PRT
          14 <213> ORGANISM: Bovine
W-->      15 <400> SEQUENCE: 1
          16 Met Lys Ala Val Gly Ala Trp Leu Leu Cys Leu Leu Leu Leu Gly Leu
          17   1               5                  10                  15
          18 Ala Leu Gln Gly Ala Ala Ser Arg Ala His Gln His Ser Met Glu Ile
          19              20                  25                  30
          20 Arg Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg
          21          35                  40                  45
          22 Pro Val Gly Arg Phe Gly Arg Arg Arg Ala Ala Pro Gly Asp Gly Pro
          23      50                  55                  60
          24 Arg Pro Gly Pro Arg Arg Val Pro Ala Cys Phe Arg Leu Glu Gly Gly
          25  65                  70                  75                  80
          26 Ala Glu Pro Ser Arg Ala Leu Pro Gly Arg Leu Thr Ala Gln Leu Val
          27              85                  90                  95
          28 Gln Glu
          31 <210> SEQ ID NO: 2
          32 <211> LENGTH: 294
          33 <212> TYPE: DNA
          34 <213> ORGANISM: Bovine
W-->      35 <400> SEQUENCE: 2
C-->      36 atgaaggcgg tggggggcctg gctcctctgc ctgctgctgc tgggcctggc cctgcagggg    60
          37 gctgccagca gagcccacca gcactccatg gagatccgca cccccgacat caaccctgcc   120
          38 tggtacgcrg gccgtgggat ccggcccgtg ggccgcttcg gccggcgaag agctgcccyg   180
          39 ggggacggac ccaggcctgg cccccggcgt gtgccggcct gcttccgcct ggaaggcggy   240
          40 gctgagccct cccgagccct cccggggcgg ctgacggccc agctggtcca ggaa          294
          43 <210> SEQ ID NO: 3
          44 <211> LENGTH: 31
          45 <212> TYPE: PRT
          46 <213> ORGANISM: Bovine
W-->      47 <400> SEQUENCE: 3
          48 Ser Arg Ala His Gln His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
          49   1               5                  10                  15
          50 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Phe
          51              20                  25                  30
          53 <210> SEQ ID NO: 4
          54 <211> LENGTH: 32
```

all bases must be in lowercase letters, when using new sequence Rules format (global error)

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/868,885

DATE: 07/09/2001
TIME: 10:11:18

Input Set : A:\2584WOOP.txt.txt
Output Set: N:\CRF3\07092001\I868885.raw

```
    55 <212> TYPE: PRT
    56 <213> ORGANISM: Bovine
W--> 57 <400> SEQUENCE: 4
    58 Ser Arg Ala His Gln His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
    59   1                   5                  10                  15
    60 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Phe Gly
    61              20                  25                  30
    63 <210> SEQ ID NO: 5
    64 <211> LENGTH: 33
    65 <212> TYPE: PRT
    66 <213> ORGANISM: Bovine
W--> 67 <400> SEQUENCE: 5
    68 Ser Arg Ala His Gln His Ser Met Glu Ile Arg Thr Pro Asp Ile Asn
    69   1                   5                  10                  15
    70 Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro Val Gly Arg Phe Gly
    71              20                  25                  30
    72 Arg
    73 33
    76 <210> SEQ ID NO: 6
    77 <211> LENGTH: 20
    78 <212> TYPE: PRT
    79 <213> ORGANISM: Bovine
W--> 80 <400> SEQUENCE: 6
    81 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro
    82   1                   5                  10                  15
    83 Val Gly Arg Phe
    84              20
    87 <210> SEQ ID NO: 7
    88 <211> LENGTH: 21
    89 <212> TYPE: PRT
    90 <213> ORGANISM: Bovine
W--> 91 <400> SEQUENCE: 7
    92 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro
    93   1                   5                  10                  15
    94 Val Gly Arg Phe Gly
    95              20
    98 <210> SEQ ID NO: 8
    99 <211> LENGTH: 22
   100 <212> TYPE: PRT
   101 <213> ORGANISM: Bovine
W--> 102 <400> SEQUENCE: 8
   103 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg Pro
   104   1                   5                  10                  15
   105 Val Gly Arg Phe Gly Arg
   106              20
   109 <210> SEQ ID NO: 9
   110 <211> LENGTH: 93
   111 <212> TYPE: DNA
   112 <213> ORGANISM: Bovine
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/868,885

DATE: 07/09/2001
TIME: 10:11:18

Input Set : A:\2584WOOP.txt.txt
Output Set: N:\CRF3\07092001\I868885.raw

```
W--> 113 <400> SEQUENCE: 9
C--> 114 agcagagccc accagcactc catggagatc cgcacccccg acatcaaccc tgcctggtac   60
     115 gcrggccgtg ggatccggcc cgtgggccgc ttc                                 93
     118 <210> SEQ ID NO: 10
     119 <211> LENGTH: 96
     120 <212> TYPE: DNA
     121 <213> ORGANISM: Bovine
W--> 122 <400> SEQUENCE: 10
C--> 123 agcagagccc accagcactc catggagatc cgcacccccg acatcaaccc tgcctggtac   60
     124 gcrggccgtg ggatccggcc cgtgggccgc ttcggc                              96
     127 <210> SEQ ID NO: 11
     128 <211> LENGTH: 99
     129 <212> TYPE: DNA
     130 <213> ORGANISM: Bovine
W--> 131 <400> SEQUENCE: 11
C--> 132 agcagagccc accagcactc catggagatc cgcacccccg acatcaaccc tgcctggtac   60
     133 gcrggccgtg ggatccggcc cgtgggccgc ttcggccgg                           99
     136 <210> SEQ ID NO: 12
     137 <211> LENGTH: 60
     138 <212> TYPE: DNA
     139 <213> ORGANISM: Bovine
W--> 140 <400> SEQUENCE: 12
C--> 141 acccccgaca tcaaccctgc ctggtacgcr ggccgtggga tccggcccgt gggccgcttc   60
     144 <210> SEQ ID NO: 13
     145 <211> LENGTH: 63
     146 <212> TYPE: DNA
     147 <213> ORGANISM: Bovine
W--> 148 <400> SEQUENCE: 13
C--> 149 acccccgaca tcaaccctgc ctggtacgcr ggccgtggga tccggcccgt gggccgcttc   60
     150 ggc                                                                  63
     153 <210> SEQ ID NO: 14
     154 <211> LENGTH: 66
     155 <212> TYPE: DNA
     156 <213> ORGANISM: Bovine
W--> 157 <400> SEQUENCE: 14
C--> 158 acccccgaca tcaaccctgc ctggtacgcr ggccgtggga tccggcccgt gggccgcttc   60
     159 ggccgg                                                               66
     162 <210> SEQ ID NO: 15
     163 <211> LENGTH: 98
     164 <212> TYPE: PRT
     165 <213> ORGANISM: Bovine
W--> 166 <400> SEQUENCE: 15
     167 Met Lys Ala Val Gly Ala Trp Leu Leu Cys Leu Leu Leu Leu Gly Leu
     168   1               5                  10                  15
     169 Ala Leu Gln Gly Ala Ala Ser Arg Ala His Gln His Ser Met Glu Ile
     170          20                  25                  30
     171 Arg Thr Pro Asp Ile Asn Pro Ala Trp Tyr Ala Gly Arg Gly Ile Arg
     172          35                  40                  45
     173 Pro Val Gly Arg Phe Gly Arg Arg Arg Ala Ala Leu Gly Asp Gly Pro
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/868,885

DATE: 07/09/2001
TIME: 10:11:18

Input Set : A:\2584WOOP.txt.txt
Output Set: N:\CRF3\07092001\I868885.raw

```
174       50                    55                    60
175 Arg Pro Gly Pro Arg Arg Val Pro Ala Cys Phe Arg Leu Glu Gly Gly
176   65                  70                    75                    80
177 Ala Glu Pro Ser Arg Ala Leu Pro Gly Arg Leu Thr Ala Gln Leu Val
178                        85                    90                    95
179 Gln Glu
182 <210> SEQ ID NO: 16
183 <211> LENGTH: 83
184 <212> TYPE: PRT
185 <213> ORGANISM: Rat
W--> 186 <400> SEQUENCE: 16
187 Met Ala Leu Lys Thr Trp Leu Leu Cys Leu Leu Leu Leu Ser Leu Val
188   1               5                    10                    15
189 Leu Pro Gly Ala Ser Ser Arg Ala His Gln His Ser Met Glu Thr Arg
190                  20                    25                    30
191 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro
192             35                    40                    45
193 Val Gly Arg Phe Gly Arg Arg Arg Ala Thr Pro Arg Asp Val Thr Gly
194       50                    55                    60
195 Leu Gly Gln Leu Ser Cys Leu Pro Leu Asp Gly Arg Thr Lys Phe Ser
196   65                  70                    75                    80
197 Gln Arg Gly
200 <210> SEQ ID NO: 17
201 <211> LENGTH: 249
202 <212> TYPE: DNA
203 <213> ORGANISM: Rat
W--> 204 <400> SEQUENCE: 17
C--> 205 atggccctga agacgtggct tctgtgcttg ctgctgctaa gcttggtcct cccaggggct   60
206 tccagccgag cccaccagca ctccatggag acaagaaccc ctgatatcaa tcctgcctgg   120
207 tacacgggcc gcgggatcag gcctgtgggc cgcttcggca ggagaagggc aaccccgagg   180
208 gatgtcactg gacttggcca actcagctgc ctcccactgg atggacgcac caagttctct   240
209 cagcgtgga                                                            249
212 <210> SEQ ID NO: 18
213 <211> LENGTH: 31
214 <212> TYPE: PRT
215 <213> ORGANISM: Rat
W--> 216 <400> SEQUENCE: 18
217 Ser Arg Ala His Gln His Ser Met Glu Thr Arg Thr Pro Asp Ile Asn
218   1               5                    10                    15
219 Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro Val Gly Arg Phe
220             20                    25                    30
223 <210> SEQ ID NO: 19
224 <211> LENGTH: 32
225 <212> TYPE: PRT
226 <213> ORGANISM: Rat
W--> 227 <400> SEQUENCE: 19
228 Ser Arg Ala His Gln His Ser Met Glu Thr Arg Thr Pro Asp Ile Asn
229   1               5                    10                    15
230 Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro Val Gly Arg Phe Gly
```

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/868,885

DATE: 07/09/2001
TIME: 10:11:18

Input Set : A:\2584WO0P.txt.txt
Output Set: N:\CRF3\07092001\I868885.raw

```
231                    20                    25                    30
234 <210> SEQ ID NO: 20
235 <211> LENGTH: 33
236 <212> TYPE: PRT
237 <213> ORGANISM: Rat
W--> 238 <400> SEQUENCE: 20
239 Ser Arg Ala His Gln His Ser Met Glu Thr Arg Thr Pro Asp Ile Asn
240 1               5                   10                  15
241 Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro Val Gly Arg Phe Gly
242              20                  25                  30
243 Arg
246 <210> SEQ ID NO: 21
247 <211> LENGTH: 20
248 <212> TYPE: PRT
249 <213> ORGANISM: Rat
W--> 250 <400> SEQUENCE: 21
251 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro
252 1               5                   10                  15
253 Val Gly Arg Phe
254              20
257 <210> SEQ ID NO: 22
258 <211> LENGTH: 21
259 <212> TYPE: PRT
260 <213> ORGANISM: Rat
W--> 261 <400> SEQUENCE: 22
262 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro
263 1               5                   10                  15
264 Val Gly Arg Phe Gly
265              20
268 <210> SEQ ID NO: 23
269 <211> LENGTH: 22
270 <212> TYPE: PRT
271 <213> ORGANISM: Rat
W--> 272 <400> SEQUENCE: 23
273 Thr Pro Asp Ile Asn Pro Ala Trp Tyr Thr Gly Arg Gly Ile Arg Pro
274 1               5                   10                  15
275 Val Gly Arg Phe Gly Arg
276              20
279 <210> SEQ ID NO: 24
280 <211> LENGTH: 93
281 <212> TYPE: DNA
282 <213> ORGANISM: Rat
W--> 283 <400> SEQUENCE: 24
C--> 284 agccgagccc accagcactc catggagaca agaacccctg atatcaatcc tgcctggtac   60
285 acgggccgcg ggatcaggcc tgtgggccgc ttc                                  93
288 <210> SEQ ID NO: 25
289 <211> LENGTH: 96
290 <212> TYPE: DNA
291 <213> ORGANISM: Rat
```

```
<210> 44
<211> 31
<212> PRT
<213> Unknown
<220>
<221>
<223> Xaa on the 3rd position means Thr or Ala, Xaa on the 5th position means
Arg or Gln, Xaa on the 10th position means Ile or Thr, Xaa on the 21st position
means Thr or Ala, Xaa on the 22nd position means Gly or Ser.
<400> 44
Ser Arg Xaa His Xaa His Ser Met Glu Xaa Arg Thr Pro Asp Ile Asn
1               5                   10                  15
Pro Ala Trp Tyr Xaa Xaa Arg Gly Ile Arg Pro Val Gly Arg Phe
            20                  25                  30
```

see item 11 on Errors Summary Sheet

```
<210> 45
<211> 20
<212> PRT
<213> Unknown
<220>
<221>
<223> Xaa on the 10th position means Thr or Ala, Xaa on the 11th position means
Gly or Ser.
<400> 45
Thr Pro Asp Ile Asn Pro Ala Trp Tyr Xaa Xaa Arg Gly Ile Arg Pro
1               5                   10                  15
Val Gly Arg Phe
            20
```

same error

Input Set : **A:\2584WOOP.txt.txt**
Output Set: **N:\CRF3\07092001\I868885.raw**

```
L:4 M:283 W: Missing Blank Line separator, <120> field identifier
L:5 M:283 W: Missing Blank Line separator, <130> field identifier
L:6 M:270 C: Current Application Number differs, Replaced Current Application No
L:6 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:8 M:283 W: Missing Blank Line separator, <160> field identifier
L:15 M:283 W: Missing Blank Line separator, <400> field identifier
L:35 M:283 W: Missing Blank Line separator, <400> field identifier
L:36 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=2
L:47 M:283 W: Missing Blank Line separator, <400> field identifier
L:57 M:283 W: Missing Blank Line separator, <400> field identifier
L:67 M:283 W: Missing Blank Line separator, <400> field identifier
L:80 M:283 W: Missing Blank Line separator, <400> field identifier
L:91 M:283 W: Missing Blank Line separator, <400> field identifier
L:102 M:283 W: Missing Blank Line separator, <400> field identifier
L:113 M:283 W: Missing Blank Line separator, <400> field identifier
L:114 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=9
L:122 M:283 W: Missing Blank Line separator, <400> field identifier
L:123 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=10
L:131 M:283 W: Missing Blank Line separator, <400> field identifier
L:132 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=11
L:140 M:283 W: Missing Blank Line separator, <400> field identifier
L:141 M:112 C: (48) String data converted to lower case,
L:148 M:283 W: Missing Blank Line separator, <400> field identifier
L:149 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=13
L:157 M:283 W: Missing Blank Line separator, <400> field identifier
L:158 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=14
L:166 M:283 W: Missing Blank Line separator, <400> field identifier
L:186 M:283 W: Missing Blank Line separator, <400> field identifier
L:204 M:283 W: Missing Blank Line separator, <400> field identifier
L:205 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=17
L:216 M:283 W: Missing Blank Line separator, <400> field identifier
L:227 M:283 W: Missing Blank Line separator, <400> field identifier
L:238 M:283 W: Missing Blank Line separator, <400> field identifier
L:250 M:283 W: Missing Blank Line separator, <400> field identifier
L:261 M:283 W: Missing Blank Line separator, <400> field identifier
L:272 M:283 W: Missing Blank Line separator, <400> field identifier
L:283 M:283 W: Missing Blank Line separator, <400> field identifier
L:284 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=24
L:292 M:283 W: Missing Blank Line separator, <400> field identifier
L:293 M:112 C: (48) String data converted to lower case,
```

VERIFICATION SUMMARY
PATENT APPLICATION: **US/09/868,885**

DATE: 07/09/2001
TIME: 10:11:19

Input Set : **A:\2584WOOP.txt.txt**
Output Set: **N:\CRF3\07092001\I868885.raw**

```
M:112 Repeated in SeqNo=25
L:301 M:283 W: Missing Blank Line separator, <400> field identifier
L:302 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=26
L:310 M:283 W: Missing Blank Line separator, <400> field identifier
L:311 M:112 C: (48) String data converted to lower case,
L:318 M:283 W: Missing Blank Line separator, <400> field identifier
L:319 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=28
L:327 M:283 W: Missing Blank Line separator, <400> field identifier
L:328 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=29
L:336 M:283 W: Missing Blank Line separator, <400> field identifier
L:355 M:283 W: Missing Blank Line separator, <400> field identifier
L:356 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=31
L:367 M:283 W: Missing Blank Line separator, <400> field identifier
L:378 M:283 W: Missing Blank Line separator, <400> field identifier
L:389 M:283 W: Missing Blank Line separator, <400> field identifier
L:401 M:283 W: Missing Blank Line separator, <400> field identifier
L:412 M:283 W: Missing Blank Line separator, <400> field identifier
L:423 M:283 W: Missing Blank Line separator, <400> field identifier
L:434 M:283 W: Missing Blank Line separator, <400> field identifier
L:435 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=38
L:443 M:283 W: Missing Blank Line separator, <400> field identifier
L:444 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=39
L:452 M:283 W: Missing Blank Line separator, <400> field identifier
L:453 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=40
L:461 M:283 W: Missing Blank Line separator, <400> field identifier
L:462 M:112 C: (48) String data converted to lower case,
L:469 M:283 W: Missing Blank Line separator, <400> field identifier
L:470 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=42
L:478 M:283 W: Missing Blank Line separator, <400> field identifier
L:479 M:112 C: (48) String data converted to lower case,
M:112 Repeated in SeqNo=43
L:487 M:283 W: Missing Blank Line separator, <220> field identifier
L:488 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:44
L:490 M:283 W: Missing Blank Line separator, <400> field identifier
L:491 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:44
L:491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:493 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:44
L:493 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44
L:501 M:283 W: Missing Blank Line separator, <220> field identifier
L:502 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:45
L:504 M:283 W: Missing Blank Line separator, <400> field identifier
```

VERIFICATION SUMMARY   DATE: 07/09/2001
PATENT APPLICATION: **US/09/868,885**   TIME: 10:11:19

Input Set : **A:\2584WOOP.txt.txt**
Output Set: **N:\CRF3\07092001\I868885.raw**

L:505 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:45
L:505 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45